=> d his (FILE 'HOME' ENTERED AT 12:58:57 ON 02 DEC 2002) INDEX 'ADISALERTS, ADISINSIGHT, ADISNEWS, AGRICOLA, ANABSTR, AQUASCI, BIOBUSINESS, BIOCOMMERCE, BIOSIS, BIOTECHABS, BIOTECHDS, BIOTECHNO, CABA, CANCERLIT, CAPLUS, CEABA-VTB, CEN, CIN, CONFSCI, CROPB, CROPU, DDFB, DDFU, DGENE, DRUGB, DRUGLAUNCH, DRUGMONOG2, ...' ENTERED AT 12:59:08 ON

SEA GLYCOSYLTRANSFERASE

02 DEC 2002

FILE ADISALERTS FILE ADISINSIGHT FILE AGRICOLA 342 FILE ANABSTR 33 FILE AQUASCI 23 FILE BIOBUSINESS 69 19 FILE BIOCOMMERCE 2419 FILE BIOSIS 422 FILE BIOTECHABS 422 FILE BIOTECHDS 2295 FILE BIOTECHNO 777 FILE CABA 555 FILE CANCERLIT 3723 FILE CAPLUS 116 FILE CEABA-VTB 1.5 FILE CEN 17 FILE CIN 87 FILE CONFSCI 3 FILE CROPU FILE DDFB 129 FILE DDFU 32 FILE DGENE 1254 129 FILE DRUGB 2 FILE DRUGNL 41 FILE DRUGU 1 FILE DRUGUPDATES 33 FILE EMBAL 3398 FILE EMBASE 1935 FILE ESBIOBASE 103 FILE FEDRIP 59 FILE FROSTI 590 FILE FSTA FILE GENBANK 1363 FILE IFIPAT 165 3171 FILE JICST-EPLUS FILE KOSMET 7 728 FILE LIFESCI FILE MEDICONF 2283 FILE MEDLINE FILE NIOSHTIC 4 FILE NTIS 10 FILE OCEAN 3 4898 FILE PASCAL FILE PHAR FILE PHARMAML 5 FILE PHIN 40 FILE PROMT 2574 FILE SCISEARCH 744 FILE TOXCENTER

> FILE USPATFULL FILE USPAT2

FILE WPIDS

808

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L1

	FILE 'PASCAL, CAPLUS, EMBASE, JICST-EPLUS, SCISEARCH, BIOSIS, BIOTECHNO,
	MEDLINE, ESBIOBASE' ENTERED AT 13:00:04 ON 02 DEC 2002
L2	380 S L1 AND (NEISSERIA OR LOS OR LIPO-OLIGOSACCHARIDE)
L3	187 S L2 AND (PURIF? OR ISOLAT? OR CHARACT?)
L4	29 S L3 AND (CDNA OR CLONE)
L5	2 S L3 AND PY<1994
L6	9 DUP REM L4 (20 DUPLICATES REMOVED)
L7	2 S L2 AND PY<1994

=> dup rem 14 PROCESSING COMPLETED FOR L4 9 DUP REM L4 (20 DUPLICATES REMOVED)

=> d 16 ibib ab 1-6

ANSWER 1 OF 9 SCISEARCH COPYRIGHT 2002 ISI (R)

ACCESSION NUMBER: 2001:18476 SCISEARCH

THE GENUINE ARTICLE: 384LF

TITLE: Detection of bacterial virulence genes by subtractive

hybridization: Identification of capsular polysaccharide

of Burkholderia pseudomallei as a major virulence

determinant

Reckseidler S L; DeShazer D; Sokol P A; Woods D E AUTHOR:

(Reprint)

Univ Calgary, Hlth Sci Ctr, Dept Microbiol & Infect Dis, CORPORATE SOURCE:

3330 Hosp Dr NW, Calgary, AB T2N 4N1, Canada (Reprint); Univ Calgary, Hlth Sci Ctr, Dept Microbiol & Infect Dis, Calgary, AB T2N 4N1, Canada; USA, Med Res Inst Infect Dis,

Bacteriol Div, Ft Detrick, MD 21702 USA

COUNTRY OF AUTHOR:

Canada; USA

INFECTION AND IMMUNITY, (JAN 2001) Vol. 69, No. 1, pp. SOURCE:

34 - 44.

Publisher: AMER SOC MICROBIOLOGY, 1752 N ST NW,

WASHINGTON, DC 20036-2904 USA.

ISSN: 0019-9567. Article; Journal

DOCUMENT TYPE:

LANGUAGE: English

REFERENCE COUNT:

ABSTRACT IS AVAILABLE IN THE ALL AND IALL FORMATS

AR Burkholderia pseudomallei, the etiologic agent of melioidosis, is responsible for a broad spectrum of illnesses in humans and animals particularly in Southeast Asia and northern Australia, where it is endemic. Burkholderia thailandensis is a nonpathogenic environmental organism closely related to B. pseudomallei. Subtractive hybridization was carried out between these two species to identify genes encoding virulence determinants in B. pseudomallei. Screening of the subtraction library revealed A-T-rich DNA sequences unique to B. pseudomallei, suggesting they may have been acquired by horizontal transfer. One of the subtraction clones, pDD1015, encoded a protein with homology to a glycosyltransferase from Pseudomonas aeruginosa. This gene was insertionally inactivated in wild-type B. pseudomallei to create SR1015. It was determined by enzyme-linked immunosorbent assay and immunoelectron microscopy that the inactivated gene was involved in the production of a major surface polysaccharide. The 50% lethal dose (LD50) for wild-type B. pseudomallei is <10 CFU; the LD50 for SR1015 was determined to be 3.5 \times 10(5) CFU, similar to that of B. thailandensis (6.8 x 10(5) CFU). DNA sequencing of the region flanking the glycosyltransferase gene revealed open reading frames similar to capsular polysaccharide genes in Haemophilus influenzae, Escherichia coli, and Neisseria meningitidis. In addition, DNA from Burkholderia mallei and Burkholderia stabilis hybridized to a glycosyltransferase fragment probe, and a capsular structure was identified on the surface of B. stabilis via immunoelectron microscopy. Thus, the combination of PCR-based subtractive hybridization, insertional inactivation, and animal virulence studies has facilitated the identification of an important virulence determinant in B. pseudomallei.

ANSWER 2 OF 9 CAPLUS COPYRIGHT 2002 ACS

2000:260548 CAPLUS ACCESSION NUMBER:

DOCUMENT NUMBER: 132:289622

Neisseria branching enzyme and gene and TITLE: method for producing .alpha.-1,6-branched

.alpha.-1,4-glucans

INVENTOR(S): PATENT ASSIGNEE(S): Buttcher, Volker; Quanz, Martin

Planttec Biotechnologie G.m.b.H. Forschung &

Entwicklung, Germany; Max-Planck-Gesellschaft Zur

Forderung Der Wissenschaften E.V.

SOURCE:

PCT Int. Appl., 115 pp.

CODEN: PIXXD2

DOCUMENT TYPE:

Patent

LANGUAGE:

German

FAMILY ACC. NUM. COUNT:

PATENT INFORMATION:

PATENT NO.

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KIND DATE
                                                APPLICATION NO. DATE
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                                                 ______
                                                WO 1999-EP7562 19991008
     WO 2000022140
                        A1
                                20000420
          W: AE, AL, AM, AT, AU, AI, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU,
               CI, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL,
               IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA,
               MD, MG, MK, MN, MW, MY, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI,
               SK, SL, TJ, TM, TR, TT, UA, UG, US, UZ, VN, YU, ZA, ZW, AM, AZ,
               BY, KG, KZ, MD, RU, TJ, TM
          RW: GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE,
               DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF,
               CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
                                            DE 1998-19846635 19981009
                         A1 20000511
     DE 19846635
     DE 19924342
                         A1 20001130
                                                DE 1999-19924342 19990527
     CA 2345904
                                20000420
                                                 CA 1999-2345904 19991008
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                                                BR 1999-15026
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                         A1
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                                                EP 1999-952542 19991008
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               IE, SI, LT, LV, FI, RO
     JP 2002527068
                                                 JP 2000-576030
                        T2
                              20020827
                                                                    19991008
     WO 2000073422
                               20001207
                                                WO 2000-EP4842
                         A1
                                                                   20000526
              AE, AG, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR,
               CU, CZ, DE, DK, DM, DZ, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU,
               ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU,
               {\tt LV}, {\tt MA}, {\tt MD}, {\tt MG}, {\tt MK}, {\tt MN}, {\tt MW}, {\tt MX}, {\tt M2}, {\tt NO}, {\tt N2}, {\tt PL}, {\tt PT}, {\tt RO}, {\tt RU}, {\tt SD},
               SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, UZ, VN, YU, ZA, ZW, AM, AZ, BY, KG, KZ, MD, RU, TJ, TM
          RW: GH, GM, KE, LS, MW, MZ, SD, SL, SZ, TZ, UG, ZW, AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE, BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG
             010989 A 20020326 BR 2000-10989 20000526
244 Al 20020403 EP 2000-938690 20000526
AT, BE, CH, DE, DK, ES, FR, GB, GR, IT, LI, LU, NL, SE, MC, PT,
     BR 2000010989
     EP 1192244
               IE, SI, LT, LV, FI, RO
                                              DE 1998-19846635 A 19981009
DE 1999-19924342 A 19990527
PRIORITY APPLN. INFO.:
                                                                W 19991008
                                              WO 1999-EP7562
                                              WO 2000-EP4842
                                                                  W 20000526
```

AB The invention relates to nucleic acid mols. which code a branching enzyme from a bacterium of the genus Neisseria, to vectors, host cells, plant cells and plants contg. such nucleic acid mols., as well as to starch which can be obtained from said plants. The invention also relates to an in-vitro method for producing .alpha.-1,6-branched .alpha.-1,4-glucans based on saccharose and an enzyme combination comprised of an amylosucrase and of a branching enzyme. In addn., the invention relates to the .alpha.-1,6-branched .alpha.-1,4-glucans which can be obtained using the method. Thus, the branching enzyme gene of N. denitrificans was cloned and sequenced. This gene was expressed in potato plants, and, along with an amylosucrase gene, in Escherichia coli. glucans produced in these transgenic organisms were characterized

RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

L6 ANSWER 3 OF 9 PASCAL COPYRIGHT 2002 INIST-CNRS. ALL RIGHTS RESERVED.

DUPLICATE

AUTHOR:

ACCESSION NUMBER: 2000-0510812 PASCAL

COPYRIGHT NOTICE: Copyright .COPYRGT. 2000 INIST-CNRS. All rights

reserved.

TITLE (IN ENGLISH): Construction and characterization of

Haemophilus ducreyi lipooligosaccharide (LOS

) mutants defective in expression of heptosyltransferase III and .beta.1,4glucosyltransferase : Identification of LOS glycoforms containing lactosamine repeats

FILIATRAULT M. J.; GIBSON B. W.; SCHILLING B.; SHUHUA

SUN; MUNSON R. S. JR; CAMPAGNARI A. A.

CORPORATE SOURCE: Department of Microbiology, Division of Infectious

Diseases, University at Buffalo, Buffalo, New York

14214, United States; Center for Microbial

Pathogenesis, University at Buffalo, Buffalo, New York 14214, United States; Department of Pharmaceutical Chemistry, University of California, San Francisco, California 94143-0446, United States; Children's Research Institute, Ohio State University, Columbus,

Ohio 43205-2696, United States; Department of

Molecular Virology, Immunology, and Medical Genetics, Ohio State University, Columbus, Ohio 43205-2696, United States; Department of Medicine, Division of Infectious Diseases, University at Buffalo, Buffalo,

New York 14214, United States

SOURCE: Infection and immunity, (2000), 68(6), 3352-3361, 45

refs.

ISSN: 0019-9567 CODEN: INFIBR

DOCUMENT TYPE: Journal
BIBLIOGRAPHIC LEVEL: Analytic
COUNTRY: United States

LANGUAGE: English

AVAILABILITY: INIST-15757, 354000082410370410

To begin to understand the role of the lipooligosaccharide (LOS) molecule in chancroid infections, we constructed mutants defective in expression of glycosyltransferase genes. Pyocin lysis and immunoscreening was used to identify a LOS mutant of Haemophilus ducreyi 35000. This mutant, HD35000R, produced a LOS molecule that lacked the monoclonal antibody 3F11 epitope and migrated with an increased mobility on sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE). Structural studies indicated that the principal LOS glycoform contains lipid A, Kdo, and two of the three core heptose residues. HD35000R was transformed with a plasmid library of H. ducreyi 35000 DNA, and a clone producing the wild-type LOS was identified. Sequence analysis of the plasmid insert revealed one open reading frame (ORF) that encodes a protein with homology to the WaaQ (heptosyltransferase III) of Escherichia coli. A second ORF had homology to the LgtF (glucosyltransferase) of Neisseria meningitidis. Individual isogenic mutants lacking expression of the putative H. ducreyi heptosyltransferase III, the putative glucosyltransferase, and both glycosyltransferases were constructed and characterized. Each mutant was complemented with the representative wild-type genes in trans to restore expression of parental LOS and confirm the function of each enzyme. Matrix-assisted laser desorption ionization mass spectrometry and SDS-PAGE analysis identified several unique LOS glycoforms containing di-, tri-, and poly-N-acetyllactosamine repeats added to the terminal region of the main LOS branch synthesized by the heptosyltransferase III mutant. These novel H. ducreyi mutants provide

important tools for studying the regulation of LOS assembly and

biosynthesis.

ANSWER 4 OF 9 CAPLUS COPYRIGHT 2002 ACS ACCESSION NUMBER:

2000:235388 CAPLUS

DOCUMENT NUMBER:

133:218340

TITLE:

SOURCE:

Cloning and characterization of the

lipooligosaccharide galactosyltransferase II gene of

DUPLICATE 2

Haemophilus ducreyi

AUTHOR(S):

Sun, Shuhua; Schilling, Birgit; Tarantino, Laurie; Tullius, Michael V.; Gibson, Bradford W.; Munson,

Robert S., Jr.

CORPORATE SOURCE:

Children's Research Institute, Department of Molecular Virology, Immunology and Medical Genetics, The Ohio State University, Columbus, OH, 43205-2696, USA

Journal of Bacteriology (2000), 182(8), 2292-2298

CODEN: JOBAAY; ISSN: 0021-9193 American Society for Microbiology

PUBLISHER: DOCUMENT TYPE:

Journal

LANGUAGE:

English

Haemophilus ducreyi is the etiol. agent of chancroid, a genital ulcer disease. The lipooligosaccharide (LOS) is considered to be a major virulence determinant and has been implicated in the adherence of H. ducreyi to keratinocytes. Strain A77, an isolate from the Paris collection, is serum sensitive, poorly adherent to fibroblasts, and deficient in microcolony formation. Structural anal. indicates that the LOS of strain A77 lacks the galactose residue found in the N-acetyllactosamine portion of the strain 35000HP LOS as well as the sialic acid substitution. From an H. ducreyi 35000HP genomic DNA library, a clone complementing the defect in A77 was identified by immunol. screening with monoclonal antibody (MAb) 3F11, a MAb which recognizes the N-acetyllactosamine portion of strain 35000HP LOS The clone contained a 4-kb insert that was sequenced. One open reading frame which encodes a protein with a mol. wt. of 33,400 was identified. This protein has homol. to glycosyltransferases of Haemophilus influenzae, Haemophilus somnus, Neisseria species, and Pasteurella haemolytica. The putative H. ducreyi glycosyltransferase gene was insertionally inactivated, and an isogenic mutant of strain 35000HP was constructed. The most complex LOS glycoform produced by the mutant has a mobility on sodium dodecyl sulfate-polyacrylamide gel identical to that of the LOS of strain A77 and lacks the 3F11-binding epitope. Structural studies confirm that the most complex glycoform of the LOS isolated from the mutant lacks the galactose residue found in the N-acetyllactosamine portion of the strain 35000HP LOS. Although previously published data suggested that the serum-sensitive phenotype of

galactosyltransferase mutation in strain A77. REFERENCE COUNT:

CORPORATE SOURCE:

THERE ARE 48 CITED REFERENCES AVAILABLE FOR THIS 48 RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

ANSWER 5 OF 9 CAPLUS COPYRIGHT 2002 ACS

DUPLICATE 3

ACCESSION NUMBER: 1998:563626 CAPLUS

DOCUMENT NUMBER: 129:271347

TITLE:

Identification of a novel gene involved in pilin

glycosylation in Neisseria meningitidis

AUTHOR (S): Jennings, Michael P.; Virji, Mumtaz; Evans, Debbie;

A77 was due to the LOS mutation, we obsd. that the complemented A77 strain retained its serum-sensitive phenotype and that the

galactosyltransferase mutant retained its serum-resistant phenotype. Thus, the serum sensitivity of strain A77 cannot be attributed to the

Foster, Virginia; Srikhanta, Yoqitha N.; Steeghs,

Liana; Van Der Ley, Peter; Moxon, E. Richard Department of Microbiology, The University of

Queensland, Brisbane, 4072, Australia

SOURCE: Molecular Microbiology (1998), 29(4), 975-984 CODEN: MOMIEE; ISSN: 0950-382X

Blackwell Science Ltd.

PUBLISHER: DOCUMENT TYPE: Journal LANGUAGE: English

The pili of Neisseria meningitidis are a key virulence factor, being major adhesins of this capsulate organism that contribute to specificity for the human host. Recently it has been reported that meningococcal pili are post-translationally modified by the addn. of an O-linked trisaccharide, Gal (.beta.1-4) Gal (.alpha.1-3) 2,4-diacetimido-2,4,6-trideoxyhexose. Using a set of random genomic sequences from N. meningitidis strain MC58, the authors have identified a novel gene homologous to a family of glycosyltransferases. A plasmid clone contg. the gene was isolated from a genomic library of N. meningitidis strain MC58 and its nucleotide sequence detd. The clone contained a complete copy of the gene, here designated pglA (pilin glycosylation). Insertional mutations were constructed in pglA in a range of meningococcal strains with well-defined lipopolysaccharide (LPS) or pilin-linked glycan structures to det. whether pglA had a role in the biosynthesis of these mols. There was no alteration in the phenotype of LPS from pglA mutant strains as judged by gel migration and the binding of monoclonal antibodies. In contrast, decreased gel migration of the pilin subunit mols. of pglA mutants was obsd., which was similar to the migration of pilins of galE mutants of same strains, supporting the notion that pglA is a glycosyltransferase involved in the biosynthesis of the pilin-linked trisaccharide structure. The pglA mutation, like the galE mutation reported previously, had no effect on pilus-mediated adhesion to human epithelial or endothelial cells. Pilin from pglA mutants were unable to bind to monospecific antisera recognizing the Gal (.beta.1-4)

pilin. REFERENCE COUNT: THERE ARE 42 CITED REFERENCES AVAILABLE FOR THIS 42 RECORD. ALL CITATIONS AVAILABLE IN THE RE FORMAT

involved in the addn. of galactose of the trisaccharide substituent of

ANSWER 6 OF 9 MEDLINE 1.6

ACCESSION NUMBER: 1998048472 MEDLINE

DOCUMENT NUMBER: 98048472 PubMed ID: 9387226

TITLE: Expression of Campylobacter hyoilei lipo-

oligosaccharide (LOS) antigens in

Gal structure, suggesting that PglA is a glycosyltransferase

Escherichia coli.

AUTHOR: Korolik V; Fry B N; Alderton M R; van der Zeijst B A; Coloe

CORPORATE SOURCE: Department of Applied Biology and Biotechnology, RMIT,

Melbourne, Australia.. rabvyk@rmitcc.xx.rmit.edu.au

MICROBIOLOGY, (1997 Nov) 143 (Pt 11) 3481-9. Journal code: 9430468. ISSN: 1350-0872. SOURCE:

ENGLAND: United Kingdom PUB. COUNTRY:

DOCUMENT TYPE: Journal; Article; (JOURNAL ARTICLE)

LANGUAGE: English

FILE SEGMENT: Priority Journals

OTHER SOURCE: GENBANK-X91081; GENBANK-X91082

ENTRY MONTH: 199802

ENTRY DATE: Entered STN: 19980224

> Last Updated on STN: 19980224 Entered Medline: 19980210

AB Campylobacter spp. are well recognized as primary pathogens in animals and in people. To isolate and define the genetic regions encoding major surface antigens of Campylobacter hyoilei, genomic DNA of the type strain of the species, RMIT-32A, was cloned into a cosmid vector, pLA2917, in Escherichia coli and the resulting genomic library was screened using antiserum raised to the parent C. hyoilei strain. Six cosmid clones were found to express a series of immunoreactive bands in the 15-25 kDa range. These bands were proteinase K-resistant and were

found in the LPS fraction of the cells, suggesting that the recombinant cosmids expressed C. hyoilei <code>lipo-oligosaccharide</code> (<code>LOS</code>) antigen(s). The minimum DNA insert size required for expression of C. hyoilei <code>LOS</code> antigen(s) in E. coli was 11.8 kb. This region was subcloned into the plasmid vector pBR322. The partial sequencing of the 11.8 kb region showed that it contains two ORFs, designated rfbF and rfbP, showing homology with the rfbF gene from Serratia marcescens and the rfbP gene from Salmonella typhimurium. Both genes are involved in LPS synthesis. The region also contained a sequence homologous to the rfaC gene of E. coli and Sal. typhimurium which is involved in core oligosaccharide synthesis.

WEST

Freeform Search

Database:	US Patents Full-Text Database US Pre-Grant Publication Full-Text Database JPO Abstracts Database EPO Abstracts Database Derwent World Patents Index IBM Technical Disclosure Bulletins							
	L1 SAME (Neisseria or lipo-oligosaccharide or LOS)							
Term:								
Display: 50 Documents in Display Format: - Starting with Number 1 Generate: Hit List Hit Count Side by Side Image								
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DATE: Monday, December 02, 2002 Printable Copy Create Case

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DB=USPT,PGPB,JPAB,EPAB,DWPI; PLUR=YES; OP=ADJ				
<u>L4</u>	L1 SAME (Neisseria or lipo-oligosaccharide or LOS or lipooligosaccharide)	23	<u>L4</u>	
<u>L3</u>	L2 SAME (purif? or isolat? or charact?)	0	<u>L3</u>	
<u>L2</u>	L1 SAME (Neisseria or lipo-oligosaccharide or LOS)	23	<u>L2</u>	
<u>L1</u>	glycosyltransferase	1182	<u>L1</u>	

END OF SEARCH HISTORY

WEST

Generate Collection

Print

Search Results - Record(s) 1 through 23 of 23 returned.

L2: Entry 1 of 23

File: PGPB

Oct 17, 2002

PGPUB-DOCUMENT-NUMBER: 20020148791

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020148791 A1

TITLE: Carbohydrate purification using ultrafiltration, reverse osmosis and

nanofiltration

PUBLICATION-DATE: October 17, 2002

INVENTOR-INFORMATION:

NAME

CITY

STATE

COUNTRY

RULE-47

DeFrees, Shawn

North Wales

PΑ

US

US-CL-CURRENT: 210/767; 536/53

Full | Title | Citation | Front | Review | Classification | Date | Reference | Sequences | Attachments | Claims | Mill | Draw Desc | Image

□ 2. Document ID: US 20020132320 A1

L2: Entry 2 of 23

File: PGPB

Sep 19, 2002

PGPUB-DOCUMENT-NUMBER: 20020132320

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020132320 A1

TITLE: Glycoconjugate synthesis using a pathway-engineered organism

PUBLICATION-DATE: September 19, 2002

INVENTOR - INFORMATION:

NAME CITY STATE COUNTRY RULE-47 Wang, Peng George Troy MΙ US Chen, Xi Norristown PΑ US Liu, Ziye MΙ US Detroit Zhang, Wei Detroit MΙ US

US-CL-CURRENT: 435/193; 435/101, 435/200, 435/320.1, 435/325

Full Title Citation Front Review Classification Date Reference Sequences Attachments Claims Phili Graw Desc Image

L2: Entry 3 of 23

File: PGPB

Sep 12, 2002

PGPUB-DOCUMENT-NUMBER: 20020127682

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020127682 A1

TITLE: Glycosyltransferases for biosynthesis of oligosaccharides, and genes encoding

them

PUBLICATION-DATE: September 12, 2002

INVENTOR - INFORMATION:

NAME

CITY

STATE

COUNTRY

RULE-47

Gotschlich, Emil C.

New York

NY

US

US-CL-CURRENT: 435/193; 435/320.1, 435/325, 435/69.1, 536/23.2

Full Title Citation Front Review Classification Date Reference Sequences Attachments

KMC Draw Desc Image

___ 4. Document ID: US 20020042369 A1

L2: Entry 4 of 23

File: PGPB

Apr 11, 2002

PGPUB-DOCUMENT-NUMBER: 20020042369

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020042369 A1

TITLE: Campylobacter glycosyltransferases for biosynthesis of gangliosides and

ganglioside mimics

PUBLICATION-DATE: April 11, 2002

INVENTOR - INFORMATION:

NAME

CITY

STATE

COUNTRY RULE-47

Gilbert, Michel

Hull

CA

Wakarchuk, Warren W.

Gloucester

CA

US-CL-CURRENT: 514/12; 435/193, 435/320.1, 435/325, 536/23.2

Full Title Citation Front Review Classification Date Reference Sequences Attachments

KMMC | Draw Desc | Image |

L2: Entry 5 of 23

File: PGPB

Mar 21, 2002

PGPUB-DOCUMENT-NUMBER: 20020034805

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020034805 A1

TITLE: FUSION PROTEINS FOR USE IN ENZYMATIC SYNTHYESIS OF OLIGOSACCHARIDES

PUBLICATION-DATE: March 21, 2002

INVENTOR-INFORMATION:

NAME

CITY

STATE COUNTRY RULE-47

GILBERT, MICHEL

HULL

CA

YOUNG, N. MARTIN

GLOUCESTER

CA

WAKARCHUK, WARREN W.

GLOUCESTER

CA

US-CL-CURRENT: 435/193; 435/183, 435/200, 435/320.1, 435/325, 536/23.2

Full Title Citation Front Review Classification Date Reference Sequences Attachments

NuitC Draw Desc Image

6. Document ID: US 20020001831 A1

L2: Entry 6 of 23

File: PGPB

Jan 3, 2002

PGPUB-DOCUMENT-NUMBER: 20020001831

PGPUB-FILING-TYPE: new

DOCUMENT-IDENTIFIER: US 20020001831 A1

TITLE: Low cost manufacture of oligosaccharides

PUBLICATION-DATE: January 3, 2002

INVENTOR-INFORMATION:

NAME

CITY

STATE

COUNTRY RULE-47

Defrees, Shawn

North Wales

PΑ

US

Johnson, Karl

Willow Grove

PΑ

US

US-CL-CURRENT: 435/101; 435/84, 536/53

Full Title Citation Front Review Classification Date Reference Sequences Attachinents

KMC Drawl Desc Image

____ 7. Document ID: US 6454946 B1

L2: Entry 7 of 23

File: USPT

Sep 24, 2002

US-PAT-NO: 6454946

DOCUMENT-IDENTIFIER: US 6454946 B1

TITLE: Carbohydrate purification using ultrafiltration, reverse osmosis and

nanofiltration

Full Title Ottation Front Review Classification Date Reference Sequences Attachments

10000 Draw Desc Image

■ 8. Document ID: US 6415234 B1

L2: Entry 8 of 23

File: USPT

Jul 2, 2002

US-PAT-NO: 6415234

DOCUMENT-IDENTIFIER: US 6415234 B1

TITLE: Designing inhibitors for glycosyltransferases

Full Title Citation Front Review Classification Date Reference Sequences Attachments

PMIC Draw Desc Image

L2: Entry 9 of 23

File: USPT

Apr 30, 2002

US-PAT-NO: 6379933

DOCUMENT-IDENTIFIER: US 6379933 B1

TITLE: Method of transferring at least two saccharide units with a polyglycosyltransferase

Full Title Ortation Front Review Classification Date Reference Sequences Attachments

NVMC Draw Desc Image

☐ 10. Document ID: US 6346422 B1

L2: Entry 10 of 23

File: USPT

Feb 12, 2002

US-PAT-NO: 6346422

DOCUMENT-IDENTIFIER: US 6346422 B1

TITLE: Method of selecting bacterial strains

Full Title Citation Front Review Classification Date Reference Sequences Attachments

PMC | Draw Desc | Image

____ 11. Document ID: US 6342382 B1

L2: Entry 11 of 23

File: USPT

Jan 29, 2002

US-PAT-NO: 6342382

DOCUMENT-IDENTIFIER: US 6342382 B1

TITLE: Glycosyltransferases for biosynthesis of oligosaccharides, and genes encoding

them

Full Title Citation Front Review Classification Date Reference Sequences Attachments

NMC Draw Desc Image

☐ 12. Document ID: US 6210933 B1

L2: Entry 12 of 23

File: USPT

Apr 3, 2001

US-PAT-NO: 6210933

DOCUMENT-IDENTIFIER: US 6210933 B1

TITLE: Recombinant .alpha.-2,3-sialyltransferases and their uses

Full Title Citation Front Review Classification Date Reference Sequences Attachments

KNMC Draw Desc Image

☐ 13. Document ID: US 6204029 B1

L2: Entry 13 of 23

File: USPT

Mar 20, 2001

US-PAT-NO: 6204029

DOCUMENT-IDENTIFIER: US 6204029 B1

TITLE: Glycosylated acceptor synthesis catalyzed by glycosyl transferase and nucleotide phosphate sugar-dependent enzyme

Full Title Citation Front Review Classification Date Reference Sequences Attachments

FiniC Draw Desc Image

1 14. Document ID: US 6127153 A

L2: Entry 14 of 23

File: USPT

Oct 3, 2000

US-PAT-NO: 6127153

DOCUMENT-IDENTIFIER: US 6127153 A

TITLE: Method of transferring at least two saccharide units with a polyglycosyltransferase, a polyglycosyltransferase and gene encoding a

polyglycosyltransferase

Full Title Citation Front Review Classification Date Reference Sequences Attachments

KMMC | Erraio Desc | Image |

☐ 15. Document ID: US 6117651 A

L2: Entry 15 of 23

File: USPT

Sep 12, 2000

US-PAT-NO: 6117651

DOCUMENT-IDENTIFIER: US 6117651 A

TITLE: Expression vectors

Full Title Citation Front Review Classification Date Reference Sequences Attachments

KMC | Draw Desc | Image |

☐ 16. Document ID: US 6096529 A

L2: Entry 16 of 23

File: USPT

Aug 1, 2000

US-PAT-NO: 6096529

DOCUMENT-IDENTIFIER: US 6096529 A

TITLE: Recombinant .alpha.-2,3-sialyltransferases and their uses

Full Title Citation Front Review Classification Date Reference Sequences Attachments

KNMC | Drawn Desc | Image

L2: Entry 17 of 23

File: USPT

Aug 31, 1999

US-PAT-NO: 5945322

DOCUMENT-IDENTIFIER: US 5945322 A

TITLE: Glycosyltransferases for biosynthesis of oligosaccharides, and genes encoding

them

Full Title Citation Front Review Classification Date Reference Sequences Attachments

NuitC | Draw Desc | Image |

☐ 18. Document ID: US 5945314 A

L2: Entry 18 of 23

File: USPT

Aug 31, 1999

US-PAT-NO: 5945314

DOCUMENT-IDENTIFIER: US 5945314 A

TITLE: Process for synthesizing oligosaccharides

Full Title Citation Front Review Classification Date Reference Sequences Attachments

MMC | Draw Desc | Image |

☐ 19. Document ID: US 5798233 A

L2: Entry 19 of 23

File: USPT

Aug 25, 1998

US-PAT-NO: 5798233

DOCUMENT-IDENTIFIER: US 5798233 A

TITLE: Glycosyltransferases for biosynthesis of oligosaccharides, and genes encoding

them

Full Title Citation Front Review Classification Date Reference Sequences Attachments

MMC | Draw Desc | Image

□ 20. Document ID: US 5705367 A

L2: Entry 20 of 23

File: USPT

Jan 6, 1998

US-PAT-NO: 5705367

DOCUMENT-IDENTIFIER: US 5705367 A

TITLE: Glycosyltransferases for biosynthesis of oligosaccharides, and genes encoding

them

Full Title Citation Front Review Classification Date Reference Sequences Attachments

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☐ 21. Document ID: US 5545553 A

L2: Entry 21 of 23

File: USPT

Aug 13, 1996

US-PAT-NO: 5545553

DOCUMENT-IDENTIFIER: US 5545553 A

TITLE: Glycosyltransferases for biosynthesis of oligosaccharides, and genes encoding

them

Full Title Citation Front Review Classification Date Reference Sequences Attachments

KMC | Draw Desc | Image

☐ 22. Document ID: AU 200215769 A WO 200248320 A2

L2: Entry 22 of 23

File: DWPI

Jun 24, 2002

DERWENT-ACC-NO: 2002-583498

DERWENT-WEEK: 200267

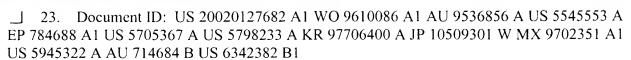
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TITLE: Novel crystal for identifying ligands that modulate glycosyltransferase activity comprises ligand binding pocket of retaining glycosyltransferase enzyme and

optionally donor and/or acceptor molecule

Full Title Citation Front Review Classification Date Reference Sequences Attachments

FNMC Draw Desc Image



L2: Entry 23 of 23

File: DWPI

Sep 12, 2002

DERWENT-ACC-NO: 1996-200924

DERWENT-WEEK: 200262

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TITLE: Nucleic acids encoding glycosyl transferase(s) - used in the diagnosis of

infection with Neisseria and for the biosynthesis of oligo:saccharide(s)

Full Title Citation Front R	eview Classification Date Reference Sequences	Attachments	KNNC Draw Desc Image
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L2: Entry 14 of 23

File: USPT

Oct 3, 2000

DOCUMENT-IDENTIFIER: US 6127153 A

TITLE: Method of transferring at least two saccharide units with a polyglycosyltransferase, a polyglycosyltransferase and gene encoding a polyglycosyltransferase

Brief Summary Text (13):

been reported as being cloned from the gonococcal strain F62 (Gotschlich, J. Exp. Dec. Med. (1994) 180, 2181-2190). Five genes lgtA, lqtB, lqtC, lqtD and lqtD. reported, and based on deletion experiments, activities are postulated, as encoding for glycosyltransferases. Due to the uncertainty caused by the nature of the deletion experiments, the exact activity of the proteins encoded by each of the genes was not ascertained and some of the genes are only suggested as being responsible for one or another activity, in the alternative. The gene lgtA is suggested as most likely to code for a GlcNAc transferase.